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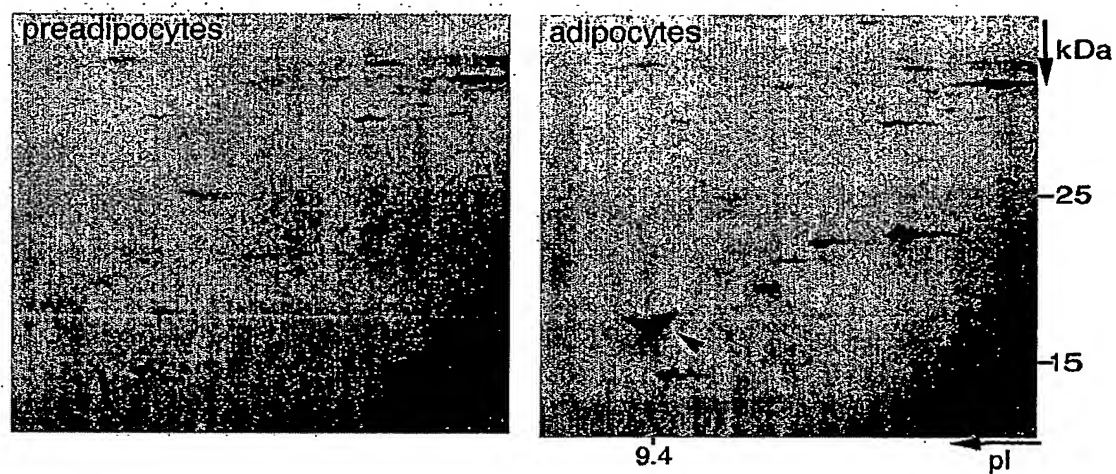
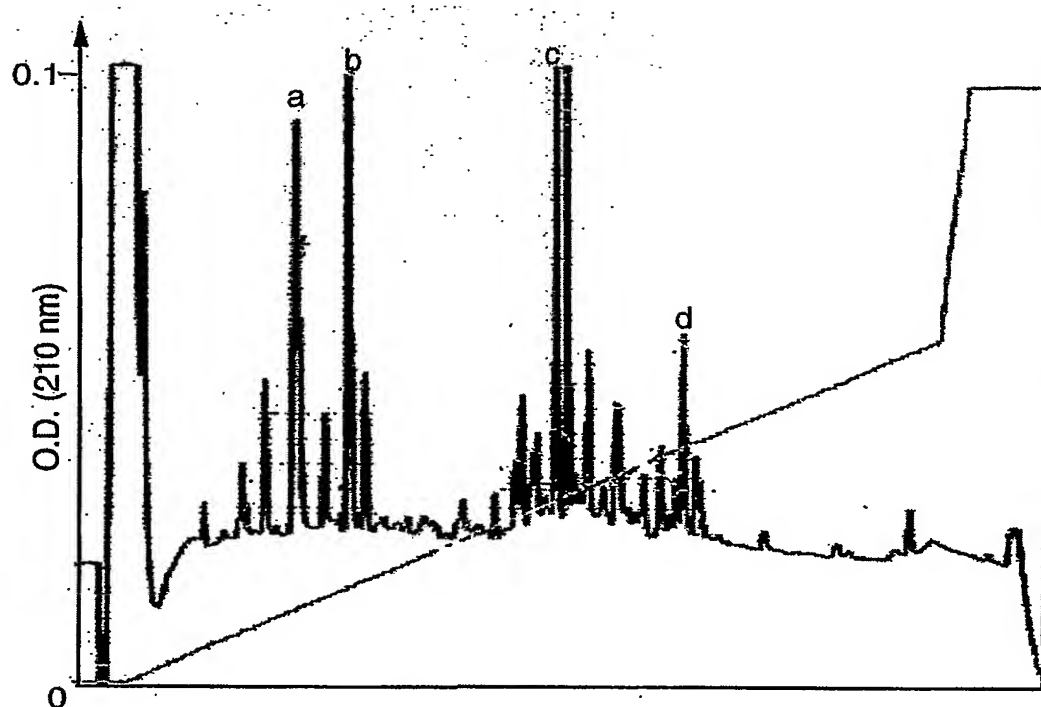


Figure 1

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Farfctions	Amino acid sequences
a	LQQTNCPK
b	GTEPELSETQR
c	CLLISLALWLGTVGTR
d	HPPVQLAFQEIGVDR

Figure 2

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Sequence ID: NO.1

MKGILSPALWECTVGRGTEPELSETQRRSLQVAL EEFHKHPPVQLA
 FQBIGVDRAEEVLFSAGTFVRLEFKLQQTNPCKDWKKPECTIKPNGR
 RRKCLACIKMDPKGKILGRIVHCPILKQGPQDPQELQCIKIAQAGEDP
 HGYFLPGQFAFSRALRTK

B.

SS (1-17)

 Potential cystatin like
 domain (32-108)

 Domain with unknown
 function (109-162)


C.

Adipocyspin

LQVLEFHHPVQLAFQEGDR EEEVLFSAGTFVR EFKLQQNPCKDWKKP-T
 90

Cystatin C

ADFVAAYNKSNDLYELRVKVS KSVV-AGNYYKVEVGEKLSPTDLNP 78

Query: 91

IKPNRRRK LACIKMD 108

Sbjct: 79

FLEQEEGFDFEVWEK 96

Figure 3